

FIGURE 1

10 20 30 40 50
MKIMMMIKLCFFSMSLICIA[↓]PADAQTEGVVFGYKGKNGPNQWGHLNPHFT
TCAVGKLQSPIDIQRRQIFYNHKLNSIHREYYFTNATLVNHVCNVAMFFG
EGAGDVIIENKNYTLLQMHWHTPSEHHLHGVQYAAELHMHQAKDGSFAV
VASLFKIGTEEPFLSQMKEKLVKLKEERLKGNHTAQVEVGRIDTRHIERK
TRKYYRYIGSLTTPPCSENSWTILGKVRSMSKEQVELLRSPLDTSFKNN
SRPCQPLNGRRVEMFHDHERVDKKETGNKKKKPN

FIGURE 2

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1  atgcagtaat ctgataaaac cctccacaga gatttccaac aaaacaggaa ctaaaacaca
61 agatgaagat tatgatgatg attaagctct gcttcttctc catgtccctc atctgcattg
121 cacctgcaga tgctcagaca gaaggagtag tgtttgata taaaggcaaa aatggacca
181 accaatgggg acacttaaac cctcacttca ccacatgcg cgtcggtaaa ttgcaatctc
241 caattgatat tcaaaggagg caaatatattt acaaccacaa attgaattca atacaccgtg
301 aatactactt cacaaacgca acactagtga accacgtctg taatgttgcc atgttcttcg
361 gggagggagc aggagatgtg ataatagaaa acaagaacta taccttactg caaatgcatt
421 ggcacactcc ttctgaacat cacctccatg gagtccaata tgcagctgag ctgcacatgg
481 tacaccaagc aaaagatgga agctttgctg tgggtggcaag tctcttcaaa atcggcactg
541 aagagccttt cctctctcag atgaaggaga aattggtgaa gctaaaggaa gagagactca
601 aagggaaacca cacagcacia gtggaagtag gaagaatcga cacaagacac attgaacgta
661 agactcgaaa gtactacaga tacattgggt cactcactac tcctccttgc tccgagaacg
721 tttcttggac catccttggc aaggtgaggt caatgtcaaa ggaacaagta gaactactca
781 gatctccatt ggacacttct ttcaagaaca attcaagacc gtgtcaaccc ctcaacggcc
841 ggagagttga gatgttccac gaccacgagc gtgtcgataa aaaagaaacc ggtaacaaaa
901 agaaaaaacc caattaaaat agttttacat tgtctattgg tttgtttaga accctaatta
961 gctttgtaaa actaataatc tcttatgtag tactgtgttg ttgtttacga cttgatatac
1021 gatttccaaa aaaaaaaaaa aaaaaa

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FIGURE 3

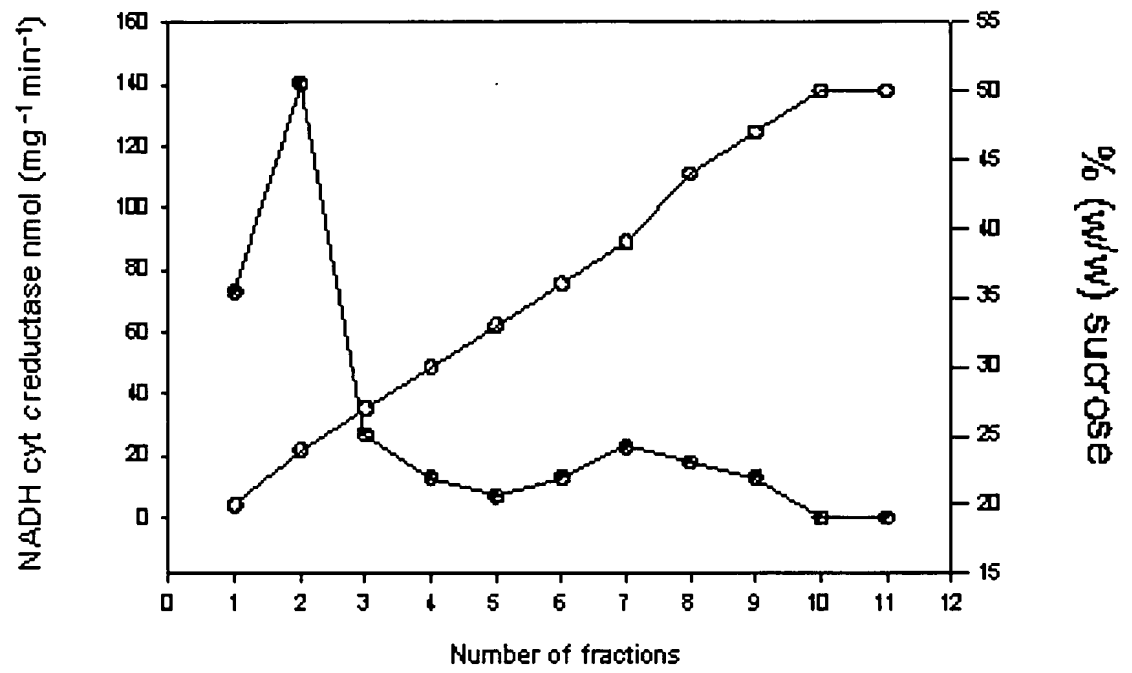


FIGURE 4

